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Sequence Listing could not be accepted.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: Thu Oct 25 18:17:00 EDT 2007
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Phe Ser Glu Pro Glu Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly
1               5                   10                  15


Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg Arg Leu Ile
                20                  25                  30


One of the above <222> lines indicates Val at locations 13 and 14; Val
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is only at location 13. Met is at location 14.

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<223> D-Ala

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Lys Lys Ile Thr Ala Gly Ala Ala Gly Val Ala Ala Gly Val Ala Ala
1   5   10   15

Ala

One of the above <223> lines indicates "D-Glu" at location 6: "Gly" is
at location 6.

****

Application No: 10583996   Version No: 2.0

Input Set:

Output Set:

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Finished: 2007-10-09 10:33:50.314
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Total Errors: 0
No. of SeqIDs Defined: 29
Actual SeqID Count: 29

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W   213   Artificial or Unknown found in <213> in SEQ ID (14)

W   213   Artificial or Unknown found in <213> in SEQ ID (15)

W   213   Artificial or Unknown found in <213> in SEQ ID (16)

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Input Set:

Output Set:

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Finished: 2007-10-09 10:33:50.314
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 618 ms
Total Warnings: 29
Total Errors: 0
No. of SeqIDs Defined: 29
Actual SeqID Count: 29

Error code   Error Description

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Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
            20   25   30

Trp Asn Trp Phe
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1   5   10   15

Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
            20   25   30

Trp Asn Trp Phe
        35

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Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly Ala Ala Gly
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Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln
            20   25   30

Leu

Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly Ala Ala Gly
1   5   10   15

Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln
            20   25   30

Leu

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Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly Ala Ala Gly
1               5                   10                  15


Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln
            20                  25                  30


Leu



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Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln
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Leu

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Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly Val Ile Gly Thr
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1   5   10   15

Thr

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Thr Lys Lys

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1   5   10   15

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1   5   10   15

Thr

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1   5   10   15

Thr Lys Lys

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1   5   10   15

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1   5   10   15

Thr

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1   5   10   15

Thr Lys Lys

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Phe Ser Glu Pro Glu Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly
1   5   10   15

Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg Arg Leu Ile
            20   25   30

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1   5   10   15

Met Ala Gly Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg
            20   25   30

Arg Leu Ile
        35

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            20   25   30

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1   5   10   15

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1               5                   10                  15


Gly Ser Leu Lys Lys
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1               5                   10                  15


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<400>  23

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1               5                   10                  15


Gly Ser Leu Lys Lys
            20



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<400>  24

Ala Val Gly Ile Gly Ala Leu Phe
1               5



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Lys Lys Ile Thr Ala Gly Ala Ala Gly Val Ala Ala Gly Val Ala Ala
1   5   10   15

Ala

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Trp Met Glu Trp Asp Arg Glu Ile Asn Asn Tyr Thr Ser Leu Ile His
1   5   10   15

Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Glu
            20   25   30

Leu Leu

<210> 27
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Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly
1   5   10

<210> 28
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Phe Phe Gly Ala Val Ile Gly Thr Ile Ala Leu Gly Val Ala Thr Ser
1   5   10   15

Ala Gln Ile Thr Ala Gly Ile Ala Leu Ala Glu Ala Arg Glu Ala Lys
            20   25   30

Arg

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<400> 29

Lys Lys Lys Met Val Leu Gly Val Phe Ala Leu Leu Phe Leu Ile Gly
1   5   10   15

Gly Ser Leu Lys Lys
            20